

5'-cca acc cag aag ccc aag gcg aat gta aat gcc-3' (SEQ ID NO:87)

5'-gtg ttc aca aca tct gcc ttg gca ttt aca atc-3' (SEQ ID NO:88)

For the mutation of lysine 15 from Trip-A:

5'-ggc att tac aat cgc ctt ggg ctt ctg ggt tgg-3' (SEQ ID NO:89)

5'-cca acc cag aag ccc aag gcg att gta aat gcc-3'--

Please amend the paragraph beginning at line 16 of page 48 with the following rewritten paragraph:

--From a cDNA library (Clontech fetal liver) the Hp-alpha sequence was PCR amplified using the following set of primers:

Non-sense primer: 5'-cac aag ctt tcc gct aga tct ctg cac tgg gtt agc cgg att ctt ggg -3' (SEQ ID NO:90)

Sense Primer: 5'-ggt gga tcc atc gag ggt agg ggt gtg gac tca ggc aat gat gtc acg g-3' -3' (SEQ ID NO:91)--

In the claims:

Please amend claims 5 and 42 as follows:

5 (Amended). The composition according to claim 2, wherein the spacer peptide comprises the amino acid sequence GTKVHMK (SEQ ID NO:69) from tetranectin, amino acid sequence PGTSGQQPSVGQQ (SEQ ID NO:70) and GTSGQ (residues 2-6 of SEQ ID